

FIG. 1

-393 TGGCCCTCGAGGCCAAGAAATTGGCACGAGGAGGGAGCCAGAGGGGGAGCCAGGACCCCTCGCTGGCTCCAGCACCCAGACCCCTGGCGG -301
-300 CGCCTCGCTTGGCAAGAGCAAGGGAGAACACTTTATATTGAACCGCGAACATTTCCTGGTCACTGAGATGAGTCTCCCGTAGGTCTGGCTTCCGCCTC -201
-200 TTATCGTGGTTGATCCCAGCTGCTCCTTCCGAACCTCCCGGGGTGCAGGCCTAGAGCCCTCCGGGGCTGACTCCAGAGTAGAGGAAGGG -101
-100 AGGGGCCCTCGGGCTGGTCCCCGAAAGCCCCCTCGCTGCCAGATGCCAGCTGGGGATGCCAGCTGGGGGGTGGCCCCGGTGGGGGAGGCCACAGCA -1
1 ATG CAG GCG CTT AAC ATT ACC CCG GAG CAG TTC TCT CGG CTG CTG CCG GAC CAC AAC CTG ACG CGG GAG CAG TTC 75
1 M Q A L N I T P E Q F S R L R D H N L T R E Q F 25
76 ATC GCT CTG TAC CGG CTC CGA CCG CTC GTC TAC ACC CCA GAG CTG CCG GGA CGC GCC AAG CTG GCC CTC GTG GTC 150
26 I A L Y R L R P L V Y T P E L P G R A K L A L V T R S K A 50
151 ACC GGC GTG CTC ATC TTC GCC CTG GCG CTC TTT GGC AAT GCT CTG GTG TTC TAC GTG GTG ACC CGC AGC AAG GCC 225
51 T G V L I F A L A L F G N A L V F Y V V T R S K A 75
226 ATG CGC ACC GTC ACC AAC ATC ATT ATC TGC TCC TTG GCG CTC AGT GAC CTG CTC ATC ACC TTC ATT CCC 300
76 M R T V T N I F I C S L A L S D L I T F F C I P 100
301 GTC ACC ATG CTC CAG AAC ATT TCC GAC AAC TGG CTG CTT GGT GCT ATT TGC AAG ATG GTG CCA ATT GTC CAG 375
101 V T M L Q N I S D N W L G G A F I C K M V P F V Q 125
376 TCT ACC GCT GPT GTG ACA GAA ATC CTC ACT ATG ACC TGC ATT GCT GTG GAA AGG CAC CAG GGA CCTT GTG CAT CCT 450
126 S T A V V T E I L T M T C I A V E R H Q G L V H P 150
451 TTT AAA ATG AAG TGG CAA TAC ACC AAC CGA AGG GCT TTC ACA ATG CTA GGT GTG GTC TGG CTG GTG GCA GTC ATC 525
151 F K M K W Q Y T N R R A F T M L G V V W L V A V I 175
526 GTA GGA TCA CCC ATG TGG CAC GTG CAA CAA CTT GAG ATC AAA TAT GAC TTC CTA TAT GAA AAG GAA CAC ATC TGC 600
176 V G S P M W H V Q Q L E I K Y D F L Y E K E H I C 200
601 TGC TTA GAA GAG TGG ACC AGC CCT GTG CAC CAG AAG ATC TAC ACC ACC TPC ATC CTT GTC ATC CTC TTC CTC CTG 675
201 C L E E W T S P V H Q K I Y T T F I L V I L F L L 225
676 CCTT ATG GTG ATG CTT ATT CTG TAC AGT AAA ATT GGT TAT GAA CTT TGG ATA AAG AAA AGA GTT GGG GAT GGT 750
226 P L M V M L I L Y S K T G Y E L W I K R V G D G 250
751 TCA GTG CTT CGA ACT ATT CAT GGA AAA GAA ATG TCC AAA ATA GCC AGG AAG AAA CGA GCT GTC ATT ATG ATG 825
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251	S	V	L	R	T	I	H	G	K	E	M	S	K	I	A	R	K	K	R	A	V	I	M	M	275	
826	GTG	ACA	GTG	GTG	GCT	CTC	TTC	TGG	GCA	CCA	TTC	CAT	GTC	CAT	GTC	CAT	ATG	ATG	ATT	GAA	TAC	AGT	ATG	900		
276	V	T	V	V	A	L	F	A	V	C	W	A	P	F	H	V	V	H	M	M	I	E	Y	S	N	300
901	TTT	GAA	AAG	GAA	TAT	GAT	GAT	GTC	ACA	ATC	AAG	ATG	ATT	TTT	GCT	ATC	GTC	CAA	ATT	ATT	GGA	TTT	TCC	AAC	TCC	975
301	F	E	K	E	Y	D	D	V	T	I	K	M	I	F	A	I	V	Q	I	I	G	F	S	N	S	325
326	I	C	N	P	I	V	Y	A	F	M	N	E	N	F	K	K	N	V	L	S	A	V	C	Y	C	350
976	ATC	TGT	AAT	CCC	ATT	GTC	TAT	GCA	TTT	ATG	AAT	GAA	AAC	TTC	AAA	AAA	AAT	GTT	TTG	TCT	GCA	GTT	TGT	TAT	TGC	1050
351	I	V	N	K	T	F	S	P	Q	R	H	G	N	S	G	I	T	M	M	R	K	K	A	K	375	
1051	ATA	GTA	AAT	AAA	ACC	TTC	TTC	TCT	CCA	GCA	CAA	AGG	CAT	GGA	AAT	TCA	GCA	ATT	ACA	ATG	CGG	AAG	AAA	GCA	AAG	1125
376	F	S	L	R	E	N	P	V	E	E	T	K	G	E	A	F	S	D	G	N	I	E	V	K	L	400
1126	T ^{TTT}	TCC	CTC	AGA	GAG	AAT	CCA	GTG	GAG	GAA	ACC	AAA	GGA	GAA	GCA	TTC	AGT	GAT	GGC	AAC	ATT	GAA	GTC	AAA	TTC	1200
401	C	E	Q	T	E	E	K	K	L	K	R	H	L	A	L	F	R	S	E	L	A	E	N	S	425	
1276	CCT	TTA	GAC	AGT	GGG	CAT	TAA	TTATAACAAATCTTCATAAT	*																	1317
426	P	L	D	S	G	H																				432

FIG. 1, cont.

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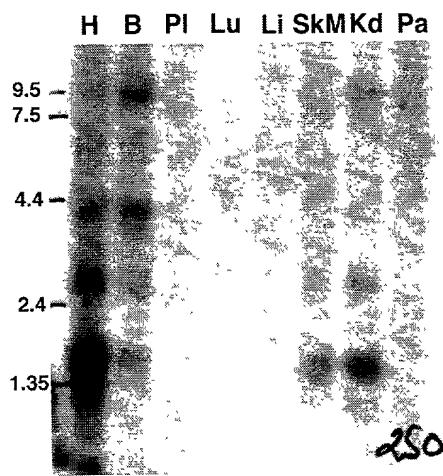


FIG. 2

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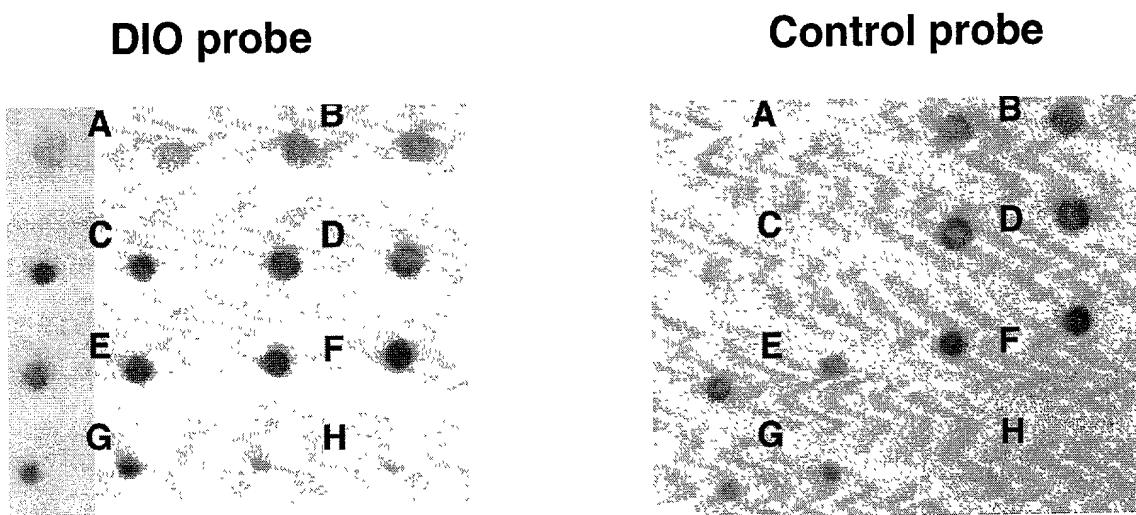


FIG. 3

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